

SEQUENCE LISTING

<110> MultiGene Biotech GmbH

<120> Novel retina-specific human proteins C7orf9, C12orf7, MPP4 and F379

<130> M36888US

<150> 60/253,751

<151> 2000-11-29

<160> 45

<170> PatentIn version 3.1

<210> 1

<211> 2435

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> artificial sequence, Translation start at 209; stop at 2435

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gttcaaggat	gaagacctac	aagagatgga	aaatttagcc	caaagaatgg	aaactcagtt	1980
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ttgatacagc gatccatagt tgcaatctaa aacaacagta ttgacccat tttaatgtgt 2220

acaactttta aagtgcagca atttattaat taatcttatt tgaaaaaaat ttttattgta 2280

tggttatgtg gttacctatt ttaacttaat ttttttctt ttacctcata tgcagctgtg 2340

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<213> Homo sapiens

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<221> misc\_feature

<223> genomic DNA, Exon from 1 to 108

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ggcaaataat cctcagttac cagaagatgt atccataact gcctagcttg cctgtcagtt 180

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Sequence file



accaggactc cttttctaga ccagaaagta atatcacctc tgacatgtga tcaaataaat 420

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cactactgcc ctaccagtga taactttaag aa 512

<210> 4

<211> 448

<212> DNA

<213> Homo sapiens

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<221> misc\_feature

<223> genomic DNA, Exon from 165 to 286

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acaaagtaag aggtggaaca gggcctgaag tcagatcttt tggcctgaga tccagtgtca 60

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acaccgcac ctgccacccc atctctgctc tcttcctttc ctaggcctct cccagatcct 180

gaggcttgat ctgcaagagc tgagtctgtt ctacagcaga gatgtgaatg gagtgtgtct 240

cttgtagat ctctccact cgccgtggct tcaggctctg ctaaaggatga gtgcttcttt 300

gctcggaagc ctttgcttgc tgaaggggtt gtggggagtg tgtagaaaat gacagcttca 360

gtccattcag gctggatagt ggaatagttt ataaacaaca gaaattgata tctcacagtt 420

Sequence

ctgtaggccca ggaagtccaa aatccagt

448

<210> 5

<211> 448

<212> DNA

<213> Homo sapiens

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<221> misc\_feature

<223> genomic DNA, Exon from 206 to 283

<400> 5

taagcttttg aagcatcggg gccaccaaac tcaagttcat ttctctttgg caactagaga 60

cacaacttac taaacaccaa ccacaccgtg ctgtgcagcc attggtgcag ttgcctgggg 120

tgtttcttct ctttgagagt cttaaatcca aaatggcaat agtcatatta tcaatatcaa 180

ttctccctcc cttgtccttc tgcagattta tgactgcctc caggaattta aagaaaagaa 240

actagttcct gccacaccac atgcacaggt gttatcctat gaggtaagga gattttattc 300

cacaggatag tagagctctg atgtggtgcc attttcccca cattgctagt tcaaatgaat 360

taaaggttct aaggaaaagt ttattgatg actatgcac taataaatgt ttctaattga 420

actttaatat aaggaagaac attggctg 448

<210> 6

<211> 384

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 165 to 245

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agatttttta aaatttttat aatgtatcct tttccatgaa ccaggtagtg gagttattac 180

gtgaaacccc tacttcccct gagatccaag agctgagaca aatgctccag gctccacact 240

tcaaggcaag tgcttgctaa aatagaaaag atgtcccat ctggcacata gacaaagttg 300

ggaaggagaa atatatgtga tggaaaatgt tctctctgaa tagatgttct attactgtac 360

acggttactg accaacagat tgta 384

<210> 7

<211> 448



<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 133 to 264

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gcgatcacat atacagtgat aataacggct gtcaactctg caagttttgc ctgtggtttc 420  
aaacatatta catgtcacg tggtttct 448

<210> 8

<211> 448

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 166 to 247

<400> 8

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tggtcatgtc aggtaacaga gggatctcgt ctattctctc ttcagggagc caccatcaag 180

cgccacgaga tgacagggga catcttggtg gccaggatca tccacggtgg gctggcggag 240

agaagtggta agctggagca gctgggattg agagttacca gaaaaacagg aaacccttga 300

ctgttttaggc ttctttctag agaaatccct tttttttctt tttttttttc tttttttttt 360

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ctgcaagctc cacctctggg gtttgcca 448

<210> 9

<211> 448

<212> DNA

<213> Homo sapiens

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<221> misc\_feature

<223> genomic DNA, Exon from 162 to 247

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ttctttgtat ctttctttgt tttttgttac tgtctgccta gggttgctat atgctggaga 180

caaactggta gaagtgaatg gagtttcagt tgagggactg gaccctgaac aagtgatcca 240

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actgggacca tcaagcccac gtgtgtgcac tgggatgtac cggggactca agttctcttg 360

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<210> 10

<211> 384

<212> DNA

<213> Homo sapiens

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<223> genomic DNA, Exon from 158 to 229

[illegible]

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<210> 14

<211> 384

<212> DNA

<213> Homo sapiens

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<221> misc\_feature

<223> genomic DNA, Exon from 179 to 217

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tctttttaat gcagacacat tttaaattct gtttctccct ttctatactc ttttatagaa 180

gatgacatga agattgatga gaaatgtgtg gaagcaggta acattttctc ttgattgctt 240

tgctgttaga agaaatatga agcatgtcaa ttatagatta tctgaagcag aggtgtccaa 300

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ctgtagttct ggcacttgga agga 384

<210> 15

<211> 320

<212> DNA

<213> Homo sapiens

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<221> misc\_feature

<223> genomic DNA, Exon from 110 to 130

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tttgaatctg gtaagtaaaa aatgagtatt tggactgat ttttaaattgt atattctaaa 180

ttttgatgca atttatacac atatttataa taactgttta aatatatcaa cattaaaaaa 240

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tttatttttta attttaattt 320

<210> 16

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<212> DNA

<213> Homo sapiens

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<221> misc\_feature

<223> genomic DNA, Exon from 174 to 188

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agtgggtgacg tgcacgtctt gcttatgtca ttgacctga tttgatggct aacatgatct 120

tcttaaaggc ttaacttttt catgtctggt tctgcactta cccaaatata cagaggaact 180

ttcagaaggt aattgttttt atttcctaga tataccaaat agaactatgt ttaagatctt 240

tcagtgcctc aaaaatgaat acttgactgg ataatgttta agatgaagat acggaatttg 300

ttgttggtta tggttttccc 320

<210> 17

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<212> DNA

<213> Homo sapiens

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<223> genomic DNA, Exon from 170 to 211

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 aaactttctcg ctctcacata ttctgctcca tctgttcttt gtgtttcaga caaggaggag 180  
 tttgttggtt acggtcagaa gttctttata ggtaggtgat aaattaacaa gaggtgggtc 240  
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<210> 18

<211> 512

<212> DNA

<213> Homo sapiens

<400> 18

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 aacacttttag gatcatgggt gctacatatt tcatcaggtg tgaagctaca agtgatctct 180  
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512

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<212> DNA

<213> Homo sapiens

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<223> genomic DNA, Exon from 160 to 240

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cacatctgat gatttctgtg tgtgactttt tgtgtttagg accctctggg gttggagtaa 180

atgagctcag aagacaactt attgaattta atcccagcca ttttcaaagt gctgtgccac 240

gtatgttttag ttctgctttc ataatggttt gtgttttggt aaaactttct ttgctgatct 300

catttaacta tgtcattcca tctttgttgt aaaagtatac aacaccaggg atagttctta 360

agtatttcta accatattta tttt 384

<210> 20

<211> 448

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 200 to 293

<400> 20

tcagtaaagg tttatagact aactgatttt gatacgagaa cttatcacca attcaggctt 60

cttcttttta gttctagcat tttatctcct tgattatata ttcatttatt tattttgatt 120

agatatcttt attcaaatgc atattggtaa tcaaagaatt ctgaagacac tgaaaccttt 180

cattcccttt ttctgataga cactactcgt actaaaaaga gttacgaaat gaatgggcgt 240

gagtatcact atgtgtccaa ggaaacattt gaaaacctca tatatagtca caggtaaagt 300

agagggttcag aagctgattc ttacctcttg ttgttttaca tttgaaatag attccctatt 360

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atgtacatcg atatacagca caccaact 448

<210> 21

<211> 448

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 133 to 241

<400> 21

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ctgctcttgc aggatgctgg agtatggtga gtacaaaggc cacctgtatg gcactagtgt 180

ggatgctgtt caaacagtcc ttgtcgaagg aaagatctgt gtcattggacc tagagcctca 240

gggtgggtcca tgggtggaata tttatgtccc caaacaatga atgcgtatca tccatttttt 300

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ggaaagaagt cttgcttctc agacagaa 448

<210> 22

<211> 448

<212> DNA

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<223> genomic DNA, Exon from 164 to 298

<400> 22

agctacttgg gaggtctgaga tgggtggatc gtttgagcct gggaagctga ggctacagtg 60

aactgtgatt gcaccacagc actccagcct ggggtgacaga gcaagaccat gtctcaaaac 120

aaaacaaaca aaaaataaat gtgcatttaa attttctgtg taggatattc aaggggttcg 180

aacctatgaa ctgaagccct atgtcatatt tataaagcca tcgaatatga ggtgtatgaa 240

acaatctcgg aaaaatgcc aaggtattac tgactactat gtggacatga agttcaagg 300

aagagcaagt caaaaactac tgtattgctt tcagtggcct ctgcgtggga gagatctggg 360

ttgggctggg ccaaggatct ctgatctcat tgcctctctc ctcttttttg accccctctc 420

caaaaggccc tcaataaaat ggtttact 448

<210> 23

<211> 704

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 197 to 704

<400> 23

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tcattctgtgc aaaattttcg gaccttactg tttttataca tagtttcaca actgaatgtg 120  
acagcataac aaactgtatt ttttccattt gtccaattaa gtctgtacta tccatatttt 180  
tctattttctc ctaaaggatg aagacctaca agagatggaa aatttagccc aaagaatgga 240  
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tgcccagttg ttgtctgcca tacagaaggc tcaggaggag cctcagtggg taccagcaac 360  
atggatttcc tcagatactg agtctcaatg agacttcttg tttaatgctg gagttttaac 420  
actgtaccct tgatacagcg atccatagtt gcaatctaaa acaacagtat ttgaccatt 480  
ttaatgtgta caactttaaa agtgcagcaa tttattaatt aatcttattt gaaaaaaatt 540  
tttattgtat ggttatgtgg ttacctattt taacttaatt ttttttcctt tacctcatat 600  
gcagctgtgg tagaaatatg aataatgtta agtcactgag tatgagaacc ttcgcagat 660  
ttcacatgat ctttttaaga tttaaataaa gagctttcct aaat 704

<210> 24

<211> 637

<212> PRT

<213> Homo sapiens

<400> 24

Met Ile Gln Ser Asp Lys Gly Ala Asp Pro Pro Asp Lys Lys Asp Met  
1 5 10 15

Lys Leu Ser Thr Ala Thr Asn Pro Gln Asn Gly Leu Ser Gln Ile Leu  
20 25 30

Arg Leu Val Leu Gln Glu Leu Ser Leu Phe Tyr Ser Arg Asp Val Asn  
35 40 45

Gly Val Cys Leu Leu Tyr Asp Leu Leu His Ser Pro Trp Leu Gln Ala  
50 55 60

Leu Leu Lys Ile Tyr Asp Cys Leu Gln Glu Phe Lys Glu Lys Lys Leu  
65 70 75 80

Val Pro Ala Thr Pro His Ala Gln Val Leu Ser Tyr Glu Val Val Glu  
85 90 95

Leu Leu Arg Glu Thr Pro Thr Ser Pro Glu Ile Gln Glu Leu Arg Gln  
100 105 110

Met Leu Gln Ala Pro His Phe Lys Ala Leu Leu Ser Ala His Asp Thr  
115 120 125



Ile Ala Gln Lys Asp Phe Glu Pro Leu Leu Pro Pro Leu Pro Asp Asn  
130 135 140

Ile Pro Glu Ser Glu Glu Ala Met Arg Ile Val Cys Leu Val Lys Asn  
145 150 155 160

Gln Gln Pro Leu Gly Ala Thr Ile Lys Arg His Glu Met Thr Gly Asp  
165 170 175

Ile Leu Val Ala Arg Ile Ile His Gly Gly Leu Ala Glu Arg Ser Gly  
180 185 190

Leu Leu Tyr Ala Gly Asp Lys Leu Val Glu Val Asn Gly Val Ser Val  
195 200 205

Glu Gly Leu Asp Pro Glu Gln Val Ile His Ile Leu Ala Met Ser Arg  
210 215 220

Gly Thr Ile Met Phe Lys Val Val Pro Val Ser Asp Pro Pro Val Asn  
225 230 235 240

Ser Gln Gln Met Val Tyr Val Arg Ala Met Thr Glu Tyr Trp Pro Gln  
245 250 255

Glu Asp Pro Asp Ile Pro Cys Met Asp Ala Gly Leu Pro Phe Gln Lys  
260 265 270

Gly Asp Ile Leu Gln Ile Val Asp Gln Asn Asp Ala Leu Trp Trp Gln  
275 280 285

Ala Arg Lys Ile Ser Asp Pro Ala Thr Cys Ala Gly Leu Val Pro Ser  
290 295 300

Asn His Leu Leu Lys Arg Lys Gln Arg Glu Phe Trp Trp Ser Gln Pro  
305 310 315 320

Tyr Gln Pro His Thr Cys Leu Lys Ser Thr Leu Ser Ile Ser Met Glu  
325 330 335

Glu Glu Asp Asp Met Lys Ile Asp Glu Lys Cys Val Glu Ala Asp Glu  
340 345 350

Glu Thr Phe Glu Ser Glu Glu Leu Ser Glu Asp Lys Glu Glu Phe Val  
355 360 365

Gly Tyr Gly Gln Lys Phe Phe Ile Ala Gly Phe Arg Arg Ser Met Arg  
370 375 380

Leu Cys Arg Arg Lys Ser His Leu Ser Pro Leu His Ala Ser Val Cys  
385 390 395 400

Cys Thr Gly Ser Cys Tyr Ser Ala Val Gly Ala Pro Tyr Glu Glu Val  
405 410 415

Val Arg Tyr Gln Arg Arg Pro Ser Asp Lys Tyr Arg Leu Ile Val Leu  
420 425 430

Met Gly Pro Ser Gly Val Gly Val Asn Glu Leu Arg Arg Gln Leu Ile  
435 440 445

Glu Phe Asn Pro Ser His Phe Gln Ser Ala Val Pro His Thr Thr Arg  
450 455 460

Thr Lys Lys Ser Tyr Glu Met Asn Gly Arg Glu Tyr His Tyr Val Ser  
465 470 475 480

Lys Glu Thr Phe Glu Asn Leu Ile Tyr Ser His Arg Met Leu Glu Tyr  
485 490 495

Gly Glu Tyr Lys Gly His Leu Tyr Gly Thr Ser Val Asp Ala Val Gln  
500 505 510

Thr Val Leu Val Glu Gly Lys Ile Cys Val Met Asp Leu Glu Pro Gln  
515 520 525

Asp Ile Gln Gly Val Arg Thr His Glu Leu Lys Pro Tyr Val Ile Phe  
530 535 540

Ile Lys Pro Ser Asn Met Arg Cys Met Lys Gln Ser Arg Lys Asn Ala  
545 550 555 560

Lys Val Ile Thr Asp Tyr Tyr Val Asp Met Lys Phe Lys Asp Glu Asp  
565 570 575

Leu Gln Glu Met Glu Asn Leu Ala Gln Arg Met Glu Thr Gln Phe Gly  
580 585 590

Gln Phe Phe Asp His Val Ile Val Asn Asp Ser Leu His Asp Ala Cys  
595 600 605

Ala Gln Leu Leu Ser Ala Ile Gln Lys Ala Gln Glu Glu Pro Gln Trp  
610 615 620

Val Pro Ala Thr Trp Ile Ser Ser Asp Thr Glu Ser Gln  
625 630 635

<210> 25

<211> 1190

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> artificial sequence, Translation start at 48, stop at 638

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<400> 25

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tttgtgcaga tgaattagtg atstccaatc ttcacagcaa agaaaattat gacaaatatt 180  
  
ctgagcctag aggataccca aaaggggaaa gaagcctcaa ttttgaggaa ttaaaagatt 240  
  
ggggacccaa aaatgttatt aagatgagta cacctgcagt caataaaatg ccacactcct 300  
  
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cagccaacct gcctctgaga tctggaagaa atatggaggt gagcctcgtg agacgtgttc 420  
  
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agaaaataga tgatgcagaa ttgaaacaag aaaaataaga aacctggagc ctgtccctaa 660  
  
agctgtggcc tgtaatctac aaatggctct atagcgaaga ccacacggaa gagtagctac 720  
  
atacacttca tcagctatgg atcatcaacg gcaatttttc cttgtcagta cagctataat 780  
  
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aaaaattaaa cagaagagca ccctgaaaaa cattatgatg gaaattaaat agtatgccag 1020  
  
aataacatgg ttgacaaata agtgaacaag gattaataat cacttacaaa cgtgtttctg 1080

tacacccttt ctatcgtgtc aaatgttaat gaatctgtga tcaattgaaa tgtaaattgtc 1140

tgtgtaaaaac tacaaaataa aaactcttag acttttagga gaaaagaaaa 1190

<210> 26

<211> 256

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 1 to 185

<400> 26

ataaacattg ggctgcacat agagacttaa ttttagattt agacaaaatg gaaattattt 60

catcaaaact attcatttta ttgacttttag ccacttcaag cttgttaaca tcaaacattt 120

tttgtgcaga tgaattagtg atstccaatc ttcacagcaa agaaaattat gacaaatatt 180

ctgaggtaag ttttttaa atctctaatg tgagtagcat taattacata atattaatcc 240

taagtcta at gatttt 256

<210> 27

<211> 512

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 62 to 462

<400> 27

gggttttaa at ctgttgctta taacaacagt atgttattgt aatggtcatt tctaattata 60

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acaaaaaat gttattaaga tgagtacacc tgcagtcaat aaaatgccac actccttcgc 180

caacttgcca ttgagatttg ggaggaacgt tcaagaagaa agaagtgctg gagcaacagc 240

caacctgcct ctgagatctg gaagaaatat ggaggtgagc ctctgtgagac gtgttcctaa 300

cctgccccaa aggtttggga gaacaacaac agccaaaagt gtctgcagga tgctgagtgga 360

tttgtgtcaa ggatccatgc attcaccatg tgccaatgac ttattttact ccatgacctg 420

ccagcaccaa gaaatccaga atcccgatca aaaacagtca aggtaaatac ctggaaacca 480

gtcaaagtgc atgggcagtt atatagaggt gg 512

<210> 28

<211> 768

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 115 to 718

<400> 28

acacaattca actcaagtat aattaggcag ttaggactat ggcttgtatt tgtatacaca 60

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gctattcaag aaaatagatg atgcagaatt gaaacaagaa aaataagaaa cctggagcct 180

gtccctaaag ctgtggcctg taatctacaa atggctctat agcgaagacc acacggaaga 240

gtagctacat acacttcacg agctatggat catcaacggc aatttttctt tgtcagtaca 300

gctataatag tatcttgaaa gttgtaaaaa aattaaagca tatttggttac gtaaagttaa 360

aatgattttt gtctgaataa aaaaaaagca ttgcaaatgc ttagaaaatc tctgataatg 420

gagagagaga cagaggaccc tcctcactac cctatataaa aatcattggc acagttacac 480

ttaataaaaa aaattaaaca gaagagcacc ctgaaaaaca ttatgatgga aattaaatag 540

tatgccagaa taacatgggt gacaaataag tgaacaagga ttaaaaaatca cttacaaacg 600

tgtttctgta caccctttct atcgtgtcaa atgttaatga atctgtgatc aattgaaatg 660

taaatgtctg tgtaaaaacta caaaataaaa actcttagac ttaggggaga aaagaaaaag 720



gcaactatga gttacctctt ttagtgtctc ctctatctac atccagaa

768

<210> 29

<211> 196

<212> PRT

<213> Homo sapiens

<400> 29

Met Glu Ile Ile Ser Ser Lys Leu Phe Ile Leu Leu Thr Leu Ala Thr  
1 5 10 15

Ser Ser Leu Leu Thr Ser Asn Ile Phe Cys Ala Asp Glu Leu Val Ile  
20 25 30

Ser Asn Leu His Ser Lys Glu Asn Tyr Asp Lys Tyr Ser Glu Pro Arg  
35 40 45

Gly Tyr Pro Lys Gly Glu Arg Ser Leu Asn Phe Glu Glu Leu Lys Asp  
50 55 60

Trp Gly Pro Lys Asn Val Ile Lys Met Ser Thr Pro Ala Val Asn Lys  
65 70 75 80

Met Pro His Ser Phe Ala Asn Leu Pro Leu Arg Phe Gly Arg Asn Val  
85 90 95

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Gln Glu Glu Arg Ser Ala Gly Ala Thr Ala Asn Leu Pro Leu Arg Ser  
100 105 110

Gly Arg Asn Met Glu Val Ser Leu Val Arg Arg Val Pro Asn Leu Pro  
115 120 125

Gln Arg Phe Gly Arg Thr Thr Thr Ala Lys Ser Val Cys Arg Met Leu  
130 135 140

Ser Asp Leu Cys Gln Gly Ser Met His Ser Pro Cys Ala Asn Asp Leu  
145 150 155 160

Phe Tyr Ser Met Thr Cys Gln His Gln Glu Ile Gln Asn Pro Asp Gln  
165 170 175

Lys Gln Ser Arg Arg Leu Leu Phe Lys Lys Ile Asp Asp Ala Glu Leu  
180 185 190

Lys Gln Glu Lys  
195

<210> 30

<211> 1188

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> artificial sequence, Translation start at 347, stop at 604

<400> 30

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gagggcaggc ctttcagctc cattctcctc caaggctgca gagggggcag gaattggggg 180

tgacaggaga gctgtaaggt ctccagtggg tcattctggg cccagagatg ggtgctgaag 240

ctcccacgcc tgctgtgaa aatggagtcc tctctcacct gggagagcca ggtgctgccc 300

cgagaaggat gcatttatgg cttcrtgaag tctttcctga cccccgatgc tgctgactat 360

agagacaaag tctcactatg ttgctcaggc tggctctgaa ctctggcct caagcgatcc 420

tcccacctya gcctcccaa gwgttgggat tatagacatg agccactgca cctggccgac 480

cttgggcaag ttcttaaacc cttcaaagcc tcatttttct ccaatcayaa aagggaaga 540

tggtaatatt ttccccwcca aattcttgtc ggatgccctc acagaattga gattatgtac 600

gtaaaacacc aggtgcctaa cccggcacag agcaggaggg ctaagcgtga catccagcac 660

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gccagactgt atttcccagc tgcccctgca gtgagatgtg gccatcggag ccagcattgg 900

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acgtgtcctc sgctgtctct tcccacytcc aaggagcacg gcaattgtgg aagaccaga 1020

ttagtgatgg cagaaccata gatgggagga acctgggtcc ctgacttaaa gtatcatgga 1080

tttgatggtt cccttagtga gaaataaact tccattgtgt ttaagccttt atttgtttat 1140

agttggttac agcaactgcc ttcttttaat taaaacactc ctgctgct 1188

<210> 31

<211> 85

<212> PRT

<213> Homo sapiens

<400> 31

Met Leu Leu Thr Ile Glu Thr Lys Ser His Tyr Val Ala Gln Ala Gly  
1 5 10 15

Leu Glu Leu Leu Ala Ser Ser Asp Pro Pro Thr Ser Ala Ser Gln Ser  
20 25 30

Val Gly Ile Ile Asp Met Ser His Cys Thr Trp Pro Thr Leu Gly Lys  
35 40 45

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Phe Leu Asn Pro Ser Lys Pro His Phe Ser Pro Ile Thr Lys Gly Lys  
50 55 60

Asp Gly Asn Ile Phe Pro Thr Lys Phe Leu Ser Asp Ala Leu Thr Glu  
65 70 75 80

Leu Arg Leu Cys Thr  
85

<210> 32

<211> 560

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 101 to 460

<400> 32

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tgttccaagg attagtcctg ctgccctgtg cccagctccc acacaacggg gtttcggggc 120

tgtggaccct gtgccaggaa aggaagggcg cagctcctgc aatgcggagc agccagggca 180

gtgggcacca ggcttttagcc tccctttctc accctacaga gggcaggccc ttcagctcca 240

ttctcctcca aggctgcaga gggggcagga attgggggtg acaggagagc tgtaaggctct	300
ccagtgggtc attctgggcc cagagatggg tgctgaagct cccacgcctg cctgtgaaaa	360
tggagtcttc tctcacctgg gagagccagg tgctgccccg agaaggatgc atttatggct	420
tcatgaagtc tttcctgacc cccgatgctg ctgactatag gtaagtctga gcaaactctgg	480
gggagcctca tcttggcatg agaaagagat ggcttcttct aagccactg gccgtgatcc	540
caggattata acacattctg	560

<210> 33

<211> 405

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 101 to 305

<400> 33

catgagagggt agtataatat agaggatatg tgtgcttact aagaggctgc ctgtctgacc	60
ttggacaagt tctttttatt tatttattta ttttttatag agacaaagtc tcaactatgtt	120
gtcagggtg gtcttgaact cctggcctca agcgatcctc ccaccttagc ctcccaaaga	180
gttgggatta tagacatgag cactgcacc tggccgacct tgggcaagtt cttaaaccct	240

Sequence file

tcaaagcctc atttttctcc aatcataaaa gggaaagatg gtaatatattt cccctccaaa 300

ttcttgtaag tattaacat tgtatatgta ttttgaacac gattaagctc taaacacttg 360

ttaggaagca ggagtagcat ttgaaacaaa cagctctttt cccac 405

<210> 34

<211> 821

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 101 to 721

<400> 34

aagtattaaa cattgtatat gtattttgaa cacgattaag ctctaaacac ttgttaggaa 60

gcaggagtag catttgaaac aaacagctct tttcccacag gtcggatgcc ctcacagaat 120

tgagattatg tacgtaaaac accaggtgcc taaccggca cagagcagga gggctaagcg 180

tgacatccag cacgtggtca gtggaatcca gtattcctac ccacctctct agtctcccct 240

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gcctgactga acattttctc cacctcctga tcatcagcag cagaaactgg ctgctcttcc 360

tcttgggtag acagccagac tgtatttccc agctgcccct gcagtgagat gtggccatcg	420
gagccagcat tggccaatgg actctgcatg ggagtgaagg atgctgcctc caggcttgtc	480
cctaaaacct cccacgtgtc ctccgcctgc tcttcccact tccaaggagc acggcaattg	540
tggaagaccc agattagtga tggcagaacc atagatggga ggaacctggg tccctgactt	600
aaagtatcat ggatttggat gttcccttag tgagaaataa acttccattg tgtttaagcc	660
tttatttgtt tatagttggt tacagcaact gccttctttt aattaaaaca ctcttgetgc	720
ttcatgttgc tggaatgctt gtaaccctgc cctgcttcac cagggttaact cctacttggc	780
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<210> 35

<211> 1514

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> artificial sequence, Translation start at 155, stop at 1192

<400> 35

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attcggagac ccagtgattg tgctccgygg agcctgggct gtgccccgcg ttgactgcct	120



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<210> 36

<211> 1544

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> artificial sequence, Translation start at 155, stop at 1222

<400> 36

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catagatacc ctacgaaccc caaatgccag ctgcatgaga aaagggactc accttctggt 180

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ggcctgtgtc	cacaatgata	ccaccagct	ccaagccata	ctggatggtg	gggtctcccc	360
agaggaggcc	accaggttg	acagcaatgg	gaggacaggc	ctcatggtcg	catgctacca	420
cggcttccag	agtgttgtgg	ccctgctcag	ccactgtcct	ttccttgatg	tgaaccagca	480
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agtgctgacc	gacagcttcg	acaccgtgtg	gaggattcgg	cagctgctga	ggcggcccca	780
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gagcctcccc	tttgccccgt	ctcctcagga	gggggggtgt	ctggaccacc	ttgtgactgc	960
cacaaccagc	ctggccagtc	ccttcgtcac	cactgcctgc	cacactctgt	gccctgacca	1020
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<210> 37

<211> 345

<212> PRT

<213> Homo sapiens

<400> 37

Met Arg Lys Gly Thr His Leu Leu Val Pro Cys Leu Glu Glu Glu Glu  
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Leu Ala Leu His Arg Arg Arg Leu Asp Met Ser Glu Ala Leu Pro Cys  
 20 25 30

Pro Gly Lys Glu Thr Pro Thr Pro Gly Cys Arg Leu Gly Ala Leu Tyr  
 35 40 45

Trp Ala Cys Val His Asn Asp Pro Thr Gln Leu Gln Ala Ile Leu Asp  
 50 55 60

Gly Gly Val Ser Pro Glu Glu Ala Thr Gln Val Asp Ser Asn Gly Arg  
65 70 75 80

Thr Gly Leu Met Val Ala Cys Tyr His Gly Phe Gln Ser Val Val Ala  
85 90 95

Leu Leu Ser His Cys Pro Phe Leu Asp Val Asn Gln Gln Asp Lys Gly  
100 105 110

Gly Asp Thr Ala Leu Met Leu Ala Ala Gln Ala Gly His Val Pro Leu  
115 120 125

Val Ser Leu Leu Leu Asn Tyr Tyr Val Gly Leu Asp Leu Glu Arg Arg  
130 135 140

Asp Gln Arg Gly Leu Thr Ala Leu Met Lys Ala Ala Met Arg Asn Arg  
145 150 155 160

Cys Ala Asp Leu Thr Ala Val Asp Pro Val Arg Gly Lys Thr Ala Leu  
165 170 175

Glu Trp Ala Val Leu Thr Asp Ser Phe Asp Thr Val Trp Arg Ile Arg  
180 185 190

Gln Leu Leu Arg Arg Pro Gln Val Glu Gln Leu Ser Gln His Tyr Lys  
195 200 205

Pro Glu Trp Pro Ala Leu Ser Gly Leu Val Ala Gln Ala Gln Ala Gln  
210 215 220

Ala Gln Val Ala Pro Ser Leu Leu Glu Arg Leu Gln Ala Thr Leu Ser  
225 230 235 240

Leu Pro Phe Ala Pro Ser Pro Gln Glu Gly Gly Val Leu Asp His Leu  
245 250 255

Val Thr Ala Thr Thr Ser Leu Ala Ser Pro Phe Val Thr Thr Ala Cys  
260 265 270

His Thr Leu Cys Pro Asp His Pro Pro Ser Leu Gly Thr Arg Ser Lys  
275 280 285

Ser Val Pro Glu Leu Leu Val Pro Ala Glu Ala Gln Ser Phe Arg Thr  
290 295 300

Pro Lys Ser Gly Pro Ser Ser Leu Ala Ile Pro Gly Ala Gln Asp Arg  
305 310 315 320

Glu Glu Glu Thr Gly Gly Gly Gly Gln Asn Gly Thr Glu Val Gly Glu  
325 330 335

Asp Gly Ile Gly Gln Ala Gly Asn Arg  
340 345

<210> 38

<211> 355

<212> PRT

<213> Homo sapiens

<400> 38

Met Arg Lys Gly Thr His Leu Leu Val Pro Cys Leu Glu Glu Glu Glu  
1 5 10 15

Leu Ala Leu His Arg Arg Arg Leu Asp Met Ser Glu Ala Leu Pro Cys  
20 25 30

Pro Gly Lys Glu Thr Pro Thr Pro Gly Cys Arg Leu Gly Ala Leu Tyr  
35 40 45

Trp Ala Cys Val His Asn Asp Pro Thr Gln Leu Gln Ala Ile Leu Asp  
50 55 60

Gly Gly Val Ser Pro Glu Glu Ala Thr Gln Val Asp Ser Asn Gly Arg  
65 70 75 80

Thr Gly Leu Met Val Ala Cys Tyr His Gly Phe Gln Ser Val Val Ala  
85 90 95

Leu Leu Ser His Cys Pro Phe Leu Asp Val Asn Gln Gln Asp Lys Gly

100

105

110

Gly Asp Thr Ala Leu Met Leu Ala Ala Gln Ala Gly His Val Pro Leu  
115 120 125

Val Ser Leu Leu Leu Asn Tyr Tyr Val Gly Leu Asp Leu Glu Arg Arg  
130 135 140

Asp Gln Arg Gly Leu Thr Ala Leu Met Lys Ala Ala Met Arg Asn Arg  
145 150 155 160

Cys Glu Cys Val Ala Thr Leu Leu Met Ala Gly Ala Asp Leu Thr Ala  
165 170 175

Val Asp Pro Val Arg Gly Lys Thr Ala Leu Glu Trp Ala Val Leu Thr  
180 185 190

Asp Ser Phe Asp Thr Val Trp Arg Ile Arg Gln Leu Leu Arg Arg Pro  
195 200 205

Gln Val Glu Gln Leu Ser Gln His Tyr Lys Pro Glu Trp Pro Ala Leu  
210 215 220

Ser Gly Leu Val Ala Gln Ala Gln Ala Gln Ala Gln Val Ala Pro Ser  
225 230 235 240

Leu Leu Glu Arg Leu Gln Ala Thr Leu Ser Leu Pro Phe Ala Pro Ser



245

250

255

Pro Gln Glu Gly Gly Val Leu Asp His Leu Val Thr Ala Thr Thr Ser

260

265

270

Leu Ala Ser Pro Phe Val Thr Thr Ala Cys His Thr Leu Cys Pro Asp

275

280

285

His Pro Pro Ser Leu Gly Thr Arg Ser Lys Ser Val Pro Glu Leu Leu

290

295

300

Val Pro Ala Glu Ala Gln Ser Phe Arg Thr Pro Lys Ser Gly Pro Ser

305

310

315

320

Ser Leu Ala Ile Pro Gly Ala Gln Asp Arg Glu Glu Glu Thr Gly Gly

325

330

335

Gly Gly Gln Asn Gly Thr Glu Val Gly Glu Asp Gly Ile Gly Gln Ala

340

345

350

Gly Asn Arg

355

&lt;210&gt; 39

&lt;211&gt; 183

&lt;212&gt; DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 1 to 143

<400> 39

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attcggagac ccagtgattg tgctccgygg agcctgggct gtgccccgcg ttgactgcct 120

catagatacc ctacgaaccc caagtaagaa aaaacgacga ccctctctcc gtgagtctca 180

ctg 183

<210> 40

<211> 462

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 108 to 358

<400> 40

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taggatccag ggagacactc actactcctc tccattctgt gttttagatg ccagctgcat 120

gagaaaaggg actcaccttc tggttccctg cctggaagag gaagagctgg cattgcacag 180

gagacggctg gacatgtctg aggcactgcc ctgcccgggc aaggagaccc ccaccccagg 240

ctgcaggctg ggggccctgt attgggcctg tgtccacaat gatcccaccc agctccaagc 300

catactggat ggtgggggtct cccagagga ggccacccag gtggacagca atgggaggg 360

gagatgtcct ggcttcccag aacagctggg ggcatctttg catccccacc acaccgtcct 420

ggcctggctc cctgagaggg gttcaggggc aatacctcct gc 462

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<210> 41

<211> 308

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 89 to 218

<400> 41

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caggggatct gagctgcccc tccctcagac aggcctcatg gtcgcatgct accacggctt 120

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ccagagtgtt gtggccctgc tcagccactg tcctttcctt gatgtgaacc agcaggacaa 180

aggaggggac acggccctca tgttggtgc ccaagcaggt gtgaggctgc tgcacccac 240

ttccgacagc ccccttttga tgcagacagg gcctcagccc cacccttggt gcacggtgtt 300

ctacacca 308

<210> 42

<211> 231

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 49 to 159

<400> 42

tcataceccc ctttcttggg gaccaagctt acccttgctg ccctgcaggc cacgtgcctc 60

tagtgagtct cctgctcaac tactatgtgg gcctggacct ggaacgccgg gaccagcggg 120

ggctcacggc gttaatgaag gctgccatgc ggaaccgctg tgagtgcgtg gccaccctcc 180

tcatggcagg tgtgcggggc ctggaccggg gtgtgtggcc tccagtcct c 231

<210> 43

<211> 231

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 49 to 189

<400> 43

tcatacccc ctttcctggg gaccaagett acccttgctg cctgcaggg cactgcctc 60

tagtgagtct cctgctcaac tactatgtgg gcctggacct ggaacgccgg gaccagcggg 120

ggctcacggc gttaatgaag gctgccatgc ggaaccgctg tgagtgcgtg gccaccctcc 180

tcatggcagg tgtgcggggc ctggaccggg gtgtgtggcc tccagtcct c 231

<210> 44

<211> 588

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 98 to 499

<400> 44

aatgtaaccc acatcagtct tgctcctaaa gaatctgccc ttccacaaat caccaacccc 60

tatcccgccc catgtcaccc cctgtgctcc ttcccaggtg ctgacctgac agcagtggac 120

cctgttcggg gcaagacggc cctggaatgg gcagtgtga ccgacagctt cgacaccgtg 180

tggaggattc ggcagctgct gaggcggccc caagtggagc agcttagcca gcaactacaag 240

cccgagtggc cggccttgtc cgggctctgt gcccaggccc aggcccaggc ccaggttgcc 300

ccttcactcc tagaacggtt gcaggctacc ttgagcctcc cctttgcccc gtctctcag 360

gaggggggtg ttctggacca ccttgtgact gccacaacca gcctggccag tcccttcgtc 420

accactgcct gccacactct gtgccctgac catccacctt cgctgggcac ccgaagcaag 480

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gggagtcccc agaggtcccc gtgggtcttc gtcccctacc agagccct 588

<210> 45

<211> 503

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<223> genomic DNA, Exon from 27 to 503

<400> 45

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gtctggccct tcctctctgg cgataccagg agctcaggat agagaagagg aaacaggagg 120

aggaggccag aatggcacag aagtagggga agatgggata ggacaggctg ggaacaggta 180

atcaggcccc tcccaggget tctttccct ctggagtgcc tccggcctcc ccatccacct 240

ctgcctaagt aaatctgctc tcaacctata tatatacaag gtcattcatt ctagcattgt 300

ttgcaagagt gaaagagtgg aaacacccga agtgtccatc agtaagggaac aggctagatt 360

gattacggat gtaattgctg tccatccata cagagcatac tctacagtgt attctaaaat 420

aagactaagg aagctgttta tattctgata tgaaactacc atcaagatgt ataaagtaaa 480

aataactaag gagtggaaca gtg 503

25

49